

SEQUENCE LISTING

<110> Alexander H. Borchers
Kenneth W. Dobie

<120> ANTISENSE MODULATION OF HEMATOPOIETIC CELL PROTEIN TYROSINE KINASE
EXPRESSION

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FOOTNOTES

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<213> Homo sapiens

 $\langle 220 \rangle$

<221> exon

 $\langle 222 \rangle \quad (4334) \dots (4437)$

<223> exon 5

<221> exon:intron junction

<222> (4437) . . . (4438)

<223> exon 5:intron 5

<221> intron

 $\langle 222 \rangle \quad (4438) \dots (8453)$

<223> intron 5

<221> intron:exon junction

<222> (8453) . . . (8454)

<223> intron 5:exon 6

<221> exon

<222> (8454) . . . (8603)

<223> exon 6

Figure 1. Schematic representation of the experimental design. The subjects were divided into two groups: the control group (CG) and the experimental group (EG). The CG was divided into two subgroups: the control group (CG) and the control group (CG). The EG was divided into two subgroups: the experimental group (EG) and the experimental group (EG). The subjects were divided into two groups: the control group (CG) and the experimental group (EG). The CG was divided into two subgroups: the control group (CG) and the control group (CG). The EG was divided into two subgroups: the experimental group (EG) and the experimental group (EG).

<223> exon 11:intron 11

<223> intron 11

<223> intron 11:exon 12

<223> exon 12

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Figure 1. Schematic representation of the experimental design. The subjects were divided into two groups: the control group (CG) and the experimental group (EG). The CG was divided into two subgroups: the control group (CG) and the control group (CG). The EG was divided into two subgroups: the experimental group (EG) and the experimental group (EG). The subjects were divided into two groups: the control group (CG) and the experimental group (EG). The CG was divided into two subgroups: the control group (CG) and the control group (CG). The EG was divided into two subgroups: the experimental group (EG) and the experimental group (EG).

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<210> 16

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<400> 36

ctgcccagct ccaagtttct

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<210> 37

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<400> 37

ggtggccatc cagacttccc

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Figure 1 consists of 12 histograms arranged in a single column. Each histogram represents the distribution of the number of non-zero elements in the vector x for a specific value of n . The x-axis for all histograms is labeled 'Number of non-zero elements' and ranges from 0 to 120. The y-axis is labeled 'Frequency' and ranges from 0 to 100. The histograms are labeled with n values: 10, 20, 30, 40, 50, 60, 70, 80, 90, 100, 110, and 120. As n increases, the distribution of non-zero elements shifts to the right, indicating that the vector x contains more non-zero elements as n increases.

<210> 38

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<400> 40

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<210> 41

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catcactttt cagaaagtcc

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<400> 48

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<400> 53

tagttcctct gctcgatgaa

20

<210> 54

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Antisense Oligonucleotide

<400> 54

tcggtggatg tagttcctct

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<210> 55

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<213> Artificial Sequence

Figure 1. The effect of the number of trials (n) on the accuracy of the estimates of the parameters of the model. The results are shown for the three models: (a) Model A, (b) Model B, and (c) Model C. The x-axis represents the number of trials (n) from 0 to 100. The y-axis represents the accuracy of the estimates from 0.0 to 1.0. The legend indicates the parameter being estimated: α (solid line), β (dashed line), and γ (dotted line). In all cases, accuracy increases with the number of trials, with α generally showing the highest accuracy and γ the lowest.

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TOGETHER WITH

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agcacactct ggatgtattc

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20

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<210> 73

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<213> Artificial Sequence

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<400> 73

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<210> 74

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Figure 1 consists of 12 sub-graphs labeled (a) through (l). Each graph plots a physiological parameter against time (0 to 10 minutes). The y-axis for all graphs ranges from 0 to 100. The x-axis for all graphs ranges from 0 to 10 minutes. The graphs show that HR, SV, CO, MAP, PVR, SVR, PPA, and PVP all increase during the intervention period, while PVP/PPA remains relatively stable.

- (a) HR (b/min): Baseline is around 70, increases to around 85 during the 10-minute period.
- (b) SV (ml): Baseline is around 70, increases to around 85 during the 10-minute period.
- (c) CO (l/min): Baseline is around 5, increases to around 7 during the 10-minute period.
- (d) MAP (mmHg): Baseline is around 70, increases to around 85 during the 10-minute period.
- (e) PVR (mmHg): Baseline is around 10, increases to around 25 during the 10-minute period.
- (f) SVR (mmHg): Baseline is around 10, increases to around 25 during the 10-minute period.
- (g) PPA (mmHg): Baseline is around 10, increases to around 25 during the 10-minute period.
- (h) PVP (mmHg): Baseline is around 10, increases to around 25 during the 10-minute period.
- (i) PVP/PPA: Baseline is around 1.0, remains relatively stable around 1.0 during the 10-minute period.
- (j) PVP/PPA: Baseline is around 1.0, remains relatively stable around 1.0 during the 10-minute period.
- (k) PVP/PPA: Baseline is around 1.0, remains relatively stable around 1.0 during the 10-minute period.
- (l) PVP/PPA: Baseline is around 1.0, remains relatively stable around 1.0 during the 10-minute period.

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<210> 83

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<212> DNA

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